



## MESP2 gene

mesoderm posterior bHLH transcription factor 2

### Normal Function

The *MESP2* gene provides instructions for making a transcription factor, which is a protein that attaches (binds) to specific regions of DNA and helps control the activity of particular genes. The MESP2 protein controls the activity of genes in the Notch pathway, an important pathway in embryonic development. The Notch pathway plays a critical role in the development of the bones of the spine (vertebrae). Specifically, the MESP2 protein and the Notch pathway are involved in separating future vertebrae from one another during early development, in a complex process called somite segmentation. Although the exact mechanism of somite segmentation is unclear, it appears to require the activity of several proteins in the Notch pathway, including the NOTCH1 protein and the MESP2 protein, to be turned on and off (oscillate) in a specific pattern.

The MESP2 protein regulates Notch activity by turning on (activating) genes in the Notch pathway, which ultimately block (repress) the activity of the NOTCH1 protein. Additionally, through unknown mechanisms, the MESP2 protein seems to mark the boundary separating future vertebrae from one another.

### Health Conditions Related to Genetic Changes

[spondylocostal dysostosis](#)

[spondylothoracic dysostosis](#)

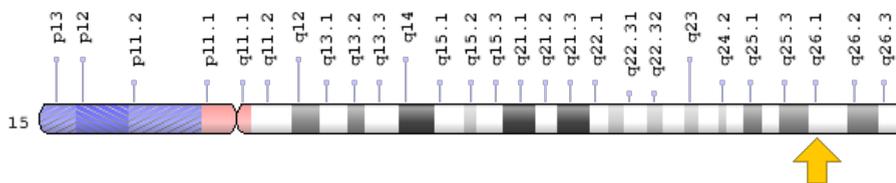
At least three mutations in the *MESP2* gene have been found to cause spondylothoracic dysostosis, a condition characterized by abnormal development of bones in the spine and ribs. All of the known mutations replace one protein building block (amino acid) in the protein sequence. The most common mutation replaces the amino acid glutamate with a premature stop signal at position 103 (written as Glu103Ter or E103X). A similar mutation occurs at amino acid position 230 (written as Glu230Ter or E230X). The third mutation replaces the amino acid leucine with the amino acid valine at position 125 (written as Leu125Val or L125V). Most affected individuals have the Glu103Ter mutation in both copies of the *MESP2* gene. However, a few people with spondylothoracic dysostosis have the Glu103Ter mutation in one copy of the *MESP2* gene and either the Leu125Val or the Glu230Ter mutation in the other copy.

Mutations in the *MESP2* gene prevent the production of any protein or lead to the production of an abnormally short, nonfunctional protein. When the MESP2 protein is nonfunctional or absent, the NOTCH1 protein is abnormally active and the boundary separating future vertebrae from one another does not form. This results in the malformation and fusion of the bones of the spine and ribs seen in spondylothoracic dysostosis.

## Chromosomal Location

Cytogenetic Location: 15q26.1, which is the long (q) arm of chromosome 15 at position 26.1

Molecular Location: base pairs 89,776,358 to 89,778,754 on chromosome 15 (Homo sapiens Annotation Release 108, GRCh38.p7) (NCBI)



Credit: Genome Decoration Page/NCBI

## Other Names for This Gene

- bHLHc6
- class C basic helix-loop-helix protein 6
- mesoderm posterior 2 homolog (mouse)
- mesoderm posterior basic helix-loop-helix transcription factor 2
- mesoderm posterior protein 2
- MESP2\_HUMAN
- SCDO2

## Additional Information & Resources

### Educational Resources

- Molecular Biology of the Cell (4th Edition, 2002): Paraxial Mesoderm: The Somites and Their Derivatives  
<https://www.ncbi.nlm.nih.gov/books/NBK10085/#A3455>

### GeneReviews

- Spondylocostal Dysostosis, Autosomal Recessive  
<https://www.ncbi.nlm.nih.gov/books/NBK8828>

### Scientific Articles on PubMed

- PubMed  
<https://www.ncbi.nlm.nih.gov/pubmed?term=%28MESP2%5BALL%5D%29+AND+english%5Bla%5D+AND+human%5Bmh%5D>

### OMIM

- MESODERM POSTERIOR bHLH TRANSCRIPTION FACTOR 2  
<http://omim.org/entry/605195>

### Research Resources

- Atlas of Genetics and Cytogenetics in Oncology and Haematology  
[http://atlasgeneticsoncology.org/Genes/GC\\_MESP2.html](http://atlasgeneticsoncology.org/Genes/GC_MESP2.html)
- ClinVar  
<https://www.ncbi.nlm.nih.gov/clinvar?term=MESP2%5Bgene%5D>
- GENATLAS  
<http://genatlas.medecine.univ-paris5.fr/fiche.php?n=29633>
- HGNC Gene Family: Basic helix-loop-helix proteins  
<http://www.genenames.org/cgi-bin/genefamilies/set/420>
- HGNC Gene Symbol Report  
[http://www.genenames.org/cgi-bin/gene\\_symbol\\_report?q=data/hgnc\\_data.php&hgnc\\_id=29659](http://www.genenames.org/cgi-bin/gene_symbol_report?q=data/hgnc_data.php&hgnc_id=29659)
- NCBI Gene  
<https://www.ncbi.nlm.nih.gov/gene/145873>
- UniProt  
<http://www.uniprot.org/uniprot/Q0VG99>

## Sources for This Summary

- Cornier AS, Staehling-Hampton K, Delventhal KM, Saga Y, Caubet JF, Sasaki N, Ellard S, Young E, Ramirez N, Carlo SE, Torres J, Emans JB, Turnpenny PD, Pourquié O. Mutations in the MESP2 gene cause spondylothoracic dysostosis/Jarcho-Levin syndrome. *Am J Hum Genet.* 2008 Jun; 82(6):1334-41. doi: 10.1016/j.ajhg.2008.04.014. Epub 2008 May 15.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/18485326>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2427230/>
- Ferjentsik Z, Hayashi S, Dale JK, Bessho Y, Herreman A, De Strooper B, del Monte G, de la Pompa JL, Maroto M. Notch is a critical component of the mouse somitogenesis oscillator and is essential for the formation of the somites. *PLoS Genet.* 2009 Sep;5(9):e1000662. doi: 10.1371/journal.pgen.1000662. Epub 2009 Sep 25.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/19779553>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2739441/>
- GeneReview: Spondylocostal Dysostosis, Autosomal Recessive  
<https://www.ncbi.nlm.nih.gov/books/NBK8828>
- Gibb S, Maroto M, Dale JK. The segmentation clock mechanism moves up a notch. *Trends Cell Biol.* 2010 Oct;20(10):593-600. doi: 10.1016/j.tcb.2010.07.001. Epub 2010 Aug 18. Review.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/20724159>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2954312/>
- Morimoto M, Takahashi Y, Endo M, Saga Y. The Mesp2 transcription factor establishes segmental borders by suppressing Notch activity. *Nature.* 2005 May 19;435(7040):354-9.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/15902259>
- Oginuma M, Takahashi Y, Kitajima S, Kiso M, Kanno J, Kimura A, Saga Y. The oscillation of Notch activation, but not its boundary, is required for somite border formation and rostral-caudal patterning within a somite. *Development.* 2010 May;137(9):1515-22. doi: 10.1242/dev.044545. Epub 2010 Mar 24.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/20335362>
- Sasaki N, Kiso M, Kitagawa M, Saga Y. The repression of Notch signaling occurs via the destabilization of mastermind-like 1 by Mesp2 and is essential for somitogenesis. *Development.* 2011 Jan;138(1):55-64. doi: 10.1242/dev.055533. Epub 2010 Nov 23.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/21098559>
- Sparrow DB, Chapman G, Turnpenny PD, Dunwoodie SL. Disruption of the somitic molecular clock causes abnormal vertebral segmentation. *Birth Defects Res C Embryo Today.* 2007 Jun;81(2): 93-110. Review.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/17600782>
- Whittock NV, Sparrow DB, Wouters MA, Sillence D, Ellard S, Dunwoodie SL, Turnpenny PD. Mutated MESP2 causes spondylocostal dysostosis in humans. *Am J Hum Genet.* 2004 Jun;74(6): 1249-54. Epub 2004 Apr 30.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/15122512>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC1182088/>

---

Reprinted from Genetics Home Reference:  
<https://ghr.nlm.nih.gov/gene/MESP2>

Reviewed: June 2016  
Published: March 21, 2017

Lister Hill National Center for Biomedical Communications  
U.S. National Library of Medicine  
National Institutes of Health  
Department of Health & Human Services